CD40T
CD40A
TNFR
FAS
ATNFR
E12
SNF1

Figure 2

Figure 1

	In-vitro- translated product		GST		GST CD40T		GST CD40A	
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Figure 3B

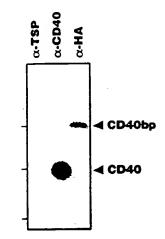


Figure 3C

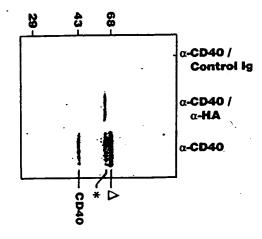
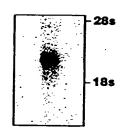


Figure 3D



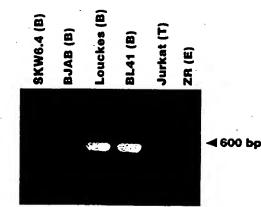


Figure 3E

165 220 275 330 385 54 495 TACQESIVKDKVFKDNÇÇKREI ELPÇVRPDÇKEKVLRKDLRDHV **X*** 기 X 3 0 FKKPTGEMNIASGCPVFVAQTVLENGTYIKD D H O 의 지 O 의 기 기 SL KGTHLSLFFVIMRGEYDALLPWPFKQKVT KTL EL **太| 団| の** SAVOHVNLLKEWSNSLEK RNNESKILHLQRVIDSQA ESVDKSAGQVARNTGLLE WKIRDYKRRKQEAVMGKTI XI KI W ഗ SVQTLLR PÇVVVSÇPHKÇS ASSAVQHVNLLK YÇRNESRĞÇAEQLMLĞHLLSSSSPKÇTAYÇRNESRÇÇYVYREATÇSHÇKSQVPMIALQKHEDTDÇPÇVV NAPSTÇSFKRYĞÇVFQGTNQQIKAHEASSA VEKNKSIQSLHNQIÇSFEIEIERQKEMLRN IRPERQNWEEADSMKSSVESLQNRVTELES LSVHDIRLADMDLGFQVLETASYNGVLIWK FVP **公**1 四1 3 ь М SAGT D.R. S S H GYKMCARVYLNGDG S DPNS NPPLKLH MLMDQGSSRRHLGDAFKP DLP PGALQT TIFIKVIVDTS SSKKMDS PFYTGY CHLVLCS LALQIYCE >l Hl 기 테 또l 있 EKAÇKYR EÇVNA OMLS 데 저 이 데 ZIOIO L S 기 이 기 포 ы

Figure 4A

igure 4B

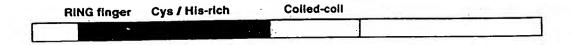
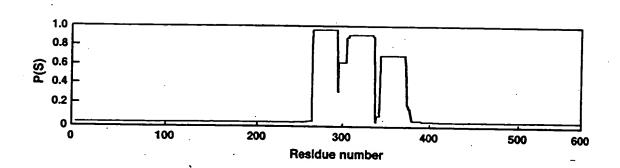


Figure 4D



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Figure .5

CACTIGGTTAAGGTCCCAGAGCTCAGAATCAGAATCAGAATCAGATCAGAAACCTGGCTCCTGGCTCCTGCTCCTGCTCCTTCTTAAGG atcecnetectralarabararcinctretiticetararigeagicgretararatesectes CCGCCGCTAAAGCTGCACACTGACCGCAGTGCTGGGACCCCAGTTTTTGTCCCTGAACAAGGAGGTTACAAGGAAAAGTTTGTGAA <u> AGAGCTGCATGGCGGCCCTGCTGAGCTCTTCAAGTCCAAAATGTACAGCGTGTCAAGAGAGGAGCATCGTTAAAGATAAGGTGTTTAAG</u> **ACAAAAGGAAATGCTTCGAAATAATGAATCCAAAATCCTTCATTTACAGCGAGTGATAGACAGCCAAGCAGAGAAACTGAAGGAG** PTGACAAGGAGGTCCGTTCCGGCAGAACTGGGAAAGGAAGCAGAAGCAGACAAGCATGAAGAGCAGCGTGGAGTCCCTCCAGAAACGGCGTC :CCGAGCTGGAGAGCGTGGACAAGAGCGCGGGGAAGTGGCTCGGAACACAGGCCTGCTGGAGTCCCAGCTGAGCCGGCATGACCA <u> ICTGGAAGATTCGCCGACTACAAGCGGCGGAAGCAGCAGCCGTCATGGGGAAQACCTGTCCCCTTTACAGCCGCTTTCTACAC</u> SCTTACTTTGCCTATAAGATGTGTCCCAGGGTCTACCTGAACGGGGACGGGATGGGGGAAGGGGACGCACTTGTCGCTGTTTTTGT TOCCCADTCTTTGTGGCCCAAACTGTTCTAGAAAATGGGACATATTAAAAGATGATACAATTTTTATTAAAGTCATAGTGGATAC ACGAAGGCCACGCCCCGGCCCCTGAGCCGGCCGAGCGGCGACGGACCGCGAGATGAGGAAAATGAGCCCCAAAGAAGTGATGATGA <u> GACCGTGGAGGACAAGTACAAGTGTGAGAAAGTGCCACCTGGTGCTGTGCAGCCCGAAGGAGGAGAGTGTGTGGGCAACTGTGCG</u> SATAATTGCTGCAAGAGAGAÄTTCTGGCTCTTCAGATCTATTGTCGGAATGAAAGCAGAGGTTGTGCAGAAGATTAATGCTGGG acctocgagaccacgtggagaaggcgtgaataccgggaagccacatgcagccactgcagagaggcaggtttccgatgatagatcs <u> BAGTGCACACTITGTCAGAGTGTGTCAATGCCCCCAGCACCTGTAGTTTTTAAGCGCTATGGCTGCGTTTTTTCAGGGGACAAACCAGC</u> ITGTTGCAGAATGAAAGTGTAGAAAAAAAAAAGAGATACAAAGTTTGCACAATCAGATATGTAGCTTTGAAAATTGAAAGT ITCGGAICTGCCCGAICCCTGATAAGTAGCTGGGAGGTGGATTTAACAGAAAGGCAACTCCTCTGGGGGATTTTGAACCGGTCTGTC CAGAAACACGAAGACACCGACTGTCCCTGCGTGGTGGTCCTGCCCTCACAAGTGCAGGCGTCCAGACTCTCCTGCTGAGGAGTGCAAATT **SAATITIATGAAATAGTA**